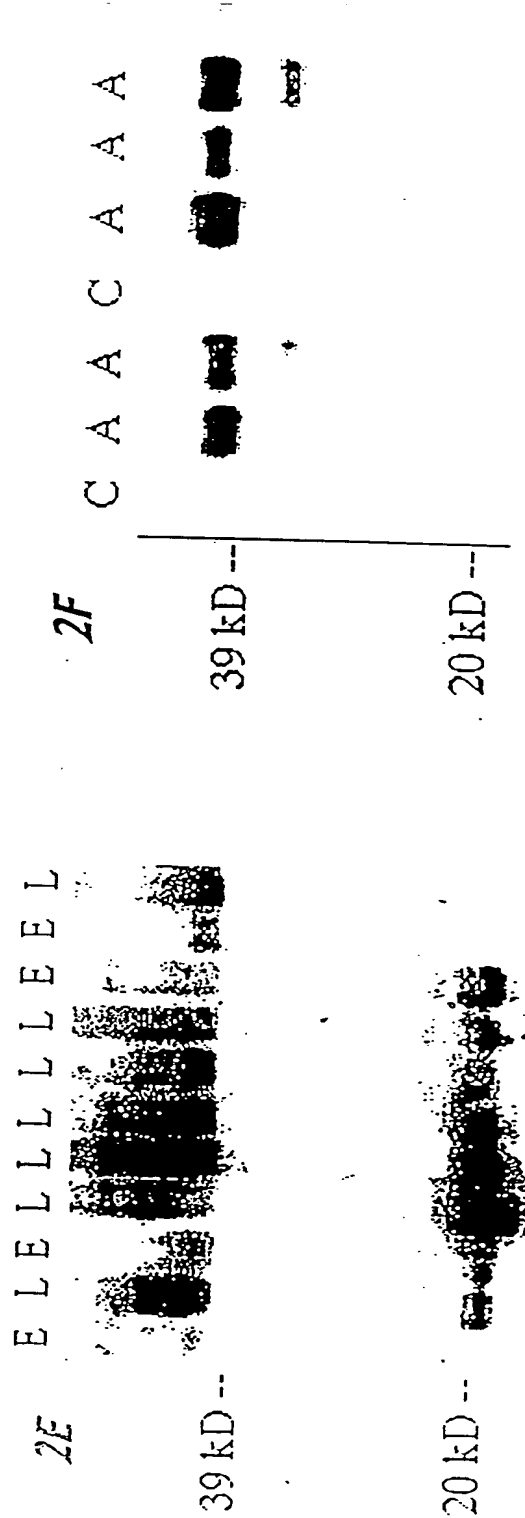
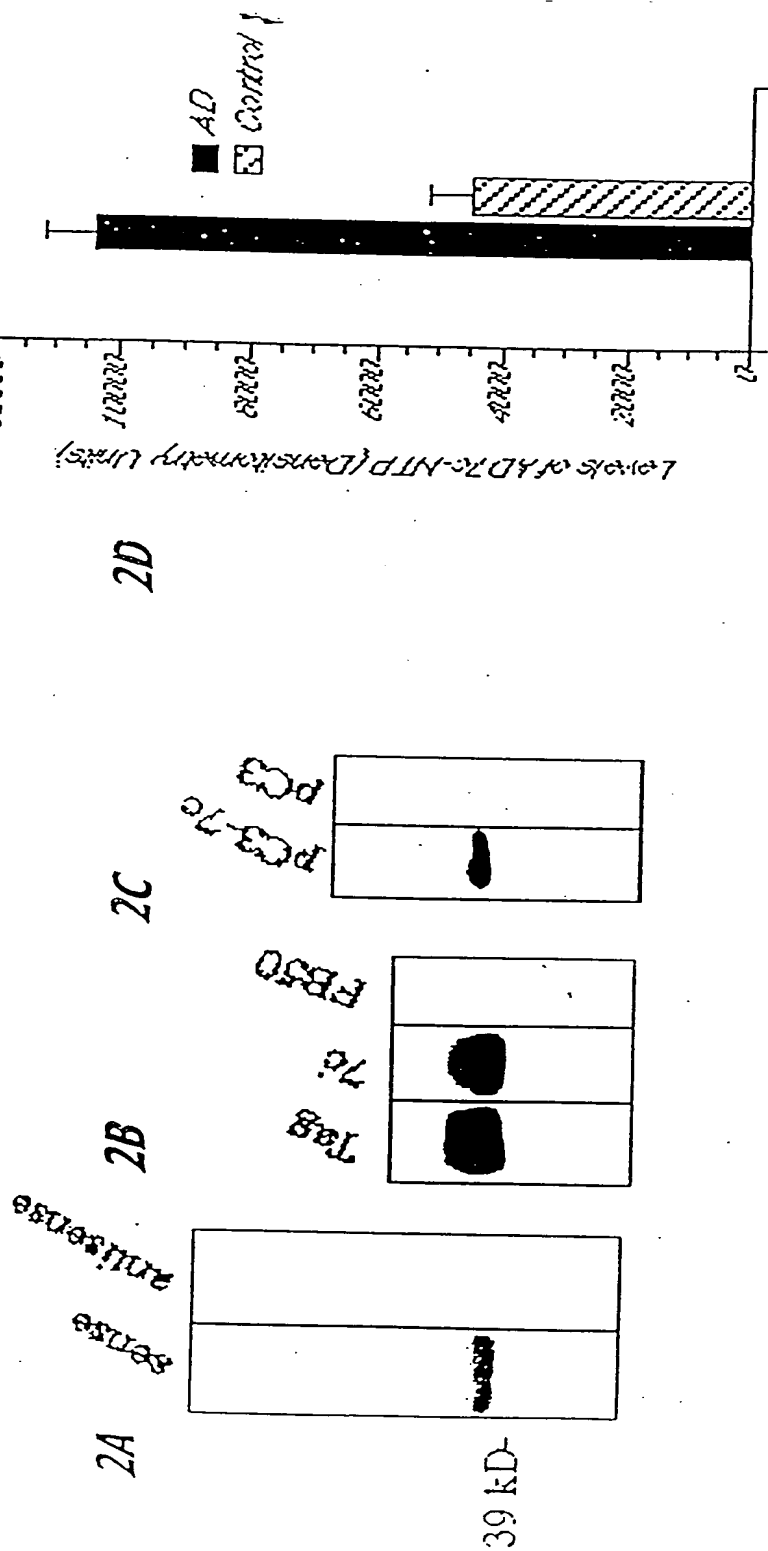
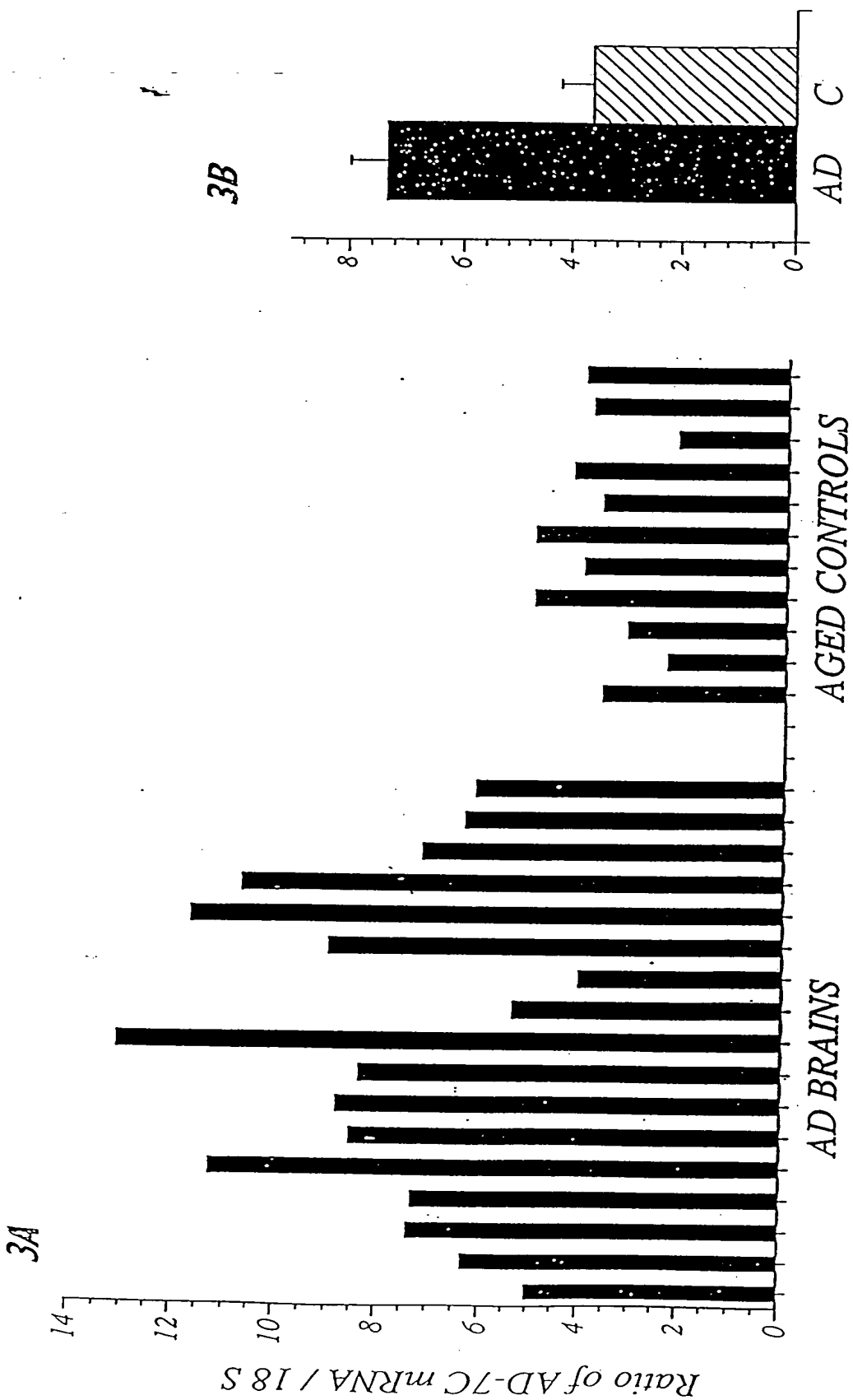


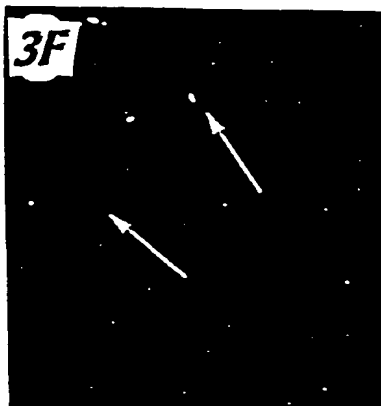
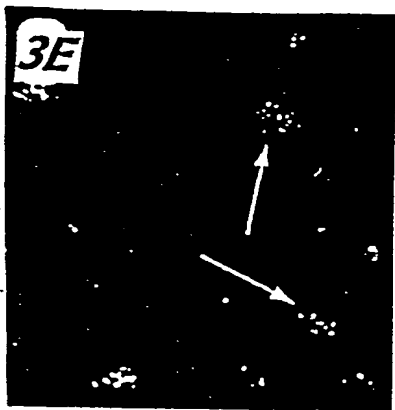
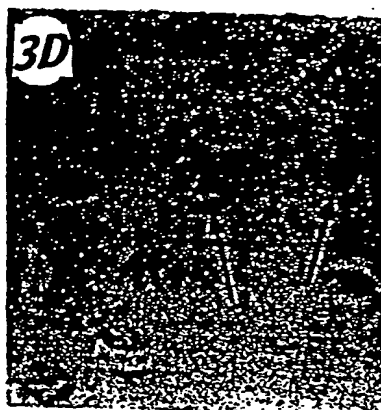
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1 M E F S L L L P R L E C N G A I 16  
63 TCA GCT CAC CGC AAC CTC CGC CTC CCG GGT TCA AGC GAT TCT CCT GCC TCA GCC TCC CCA 122  
17 S A H R N L R L P G S S D S P A S A S P 36  
123 GTA GCT GGG ATT ACA GGC ATG TGC ACC CAC GCT CGG CTA ATT TTG TAT TTT TTT TTA GTA 182  
37 V A G I T G M C T H A R L I L Y F F L V 56  
183 GAG ATG GAG TTT CTC CAT GTT GGT CAG GCT GGT CTC GAA CTC CCG ACC TCA GAT GAT CCC 242  
57 E M E F L H V G Q A G L E L P T S D D P 76  
243 TCC GTC TCG GCC TCC CAA AGT GCT AGA TAC AGG ACT GGC CAC CAT GCC CGG CTC TGC CTG 302  
77 S V S A S Q S A R Y R T G H H A R L C L 96  
303 GCT AAT TTT TGT GGT AGA AAC AGG GTT TCA CTG ATG TGC CCA AGC TGG TCT CCT GAG CTC 362  
97 A N F C G R N R V S L M C P S W S P E L 116  
363 AAG CAG TCC ACC TGC CTC AGC CTC CCA AAG TGC TGG GAT TAC AGG CGT GCA GCC GTG CCT 422  
117 K Q S T C L S L P K C W D Y R R A A V P 136  
423 GGC CTT TTT ATT TTA TTT TTT TTA AGA CAC AGG TGT GGC ACT GTC AGC GAC GAT GCA GTC 482  
137 G L F I L F F L R H R C P T L T O D E V 156  
483 GAG GAG GTC TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT 542  
157 Q W C D H S S L Q P S T P E I K H P P A 176  
543 GAG GAG GTC TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT 602  
177 S A S Q V A G T K D M H H Y T W L I F I 196  
603 GAG GAG GTC TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT 662  
197 F I F N F L R Q S L N S V T Q A G V Q W 216  
663 GAG GAG GTC TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT 722  
217 R N L G S L Q P L P P G F K L F S C P S 236  
723 GAG GAG GTC TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT 782  
237 L L S S W D Y R R P P R L A N F F V F L 256  
783 GAG GAG GTC TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT 842  
257 V E M G F T M F A R L I L I S G P C D L 276  
843 GAG GAG GTC TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT TAT 902  
277 P A S A S Q S A G I T G V S H H A R L I 296  
903 TTT AAT TTT TGT TTG TTT GAA ATG GAA TCT CAC TCT GTT ACC CAG GCT GGA GTG CAA TGG 962  
297 F N F C L F E M E S H S V T Q A G V Q W 316  
963 CCA AAT CTC GGC TCA CTG CAA CCT CTG CCT CCC GGG CTC AAG CGA TTC TCC TGT CTC AGC 1022  
317 P N L G S L Q P L P P G L K R F S C L S 336  
1023 CTC CCA AGC AGC TGG GAT TAC GGG CAC CTG CCA CCA CAC CCC GCT AAT TTT TGT ATT TTC 1082  
337 L P S S W D Y G H L P P H P A N F C I F 356  
1083 ATT AGA GGC GGG GTT TCA CCA TAT TTG TCA GGC TGG TCT CAA ACT CCT GAC CTC AGG tgac 1143  
357 I R G G V S P Y L S G W S Q T P D L R 375  
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FIG. 1

FIGS. 2A-2F 99949660

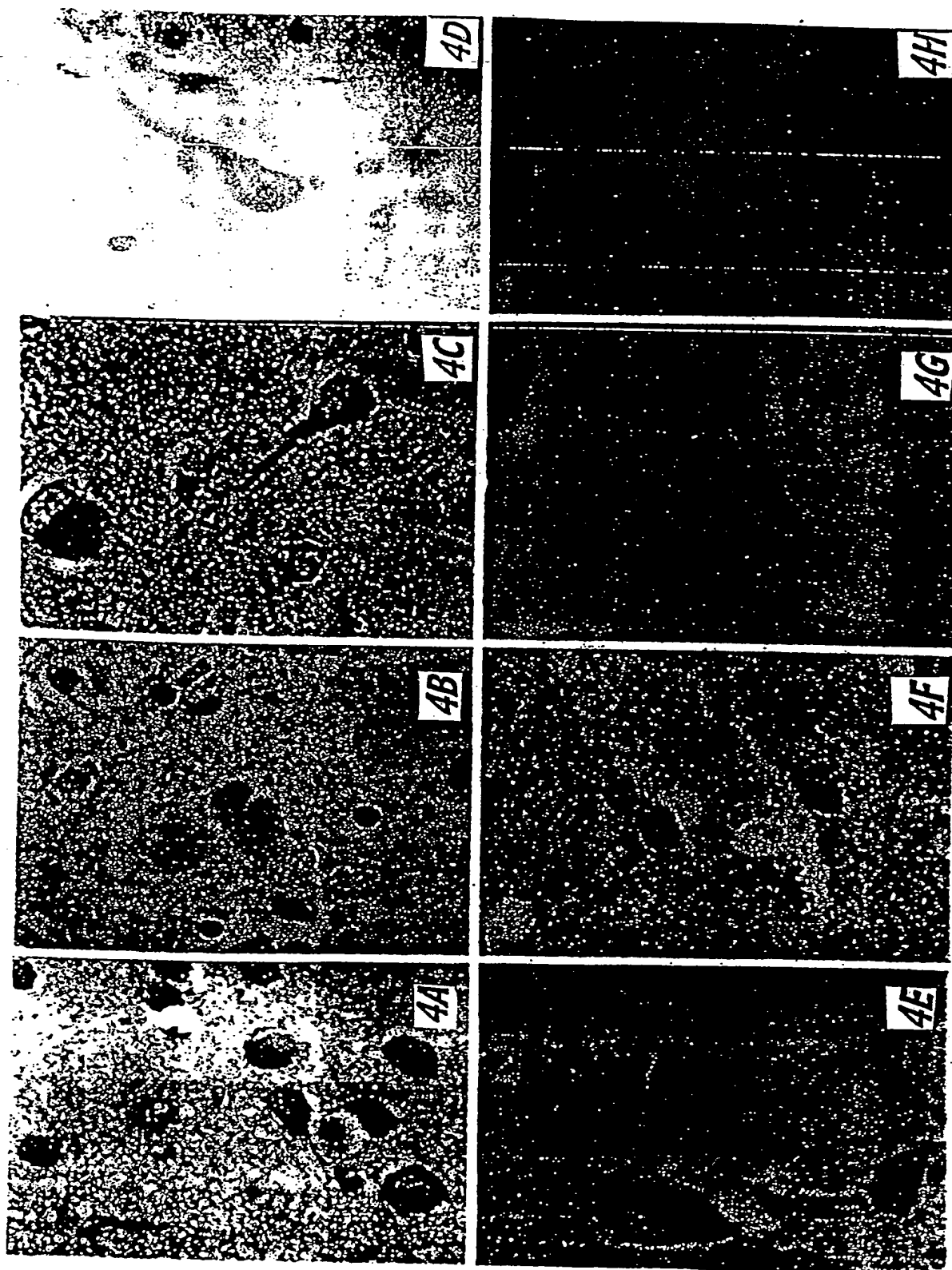




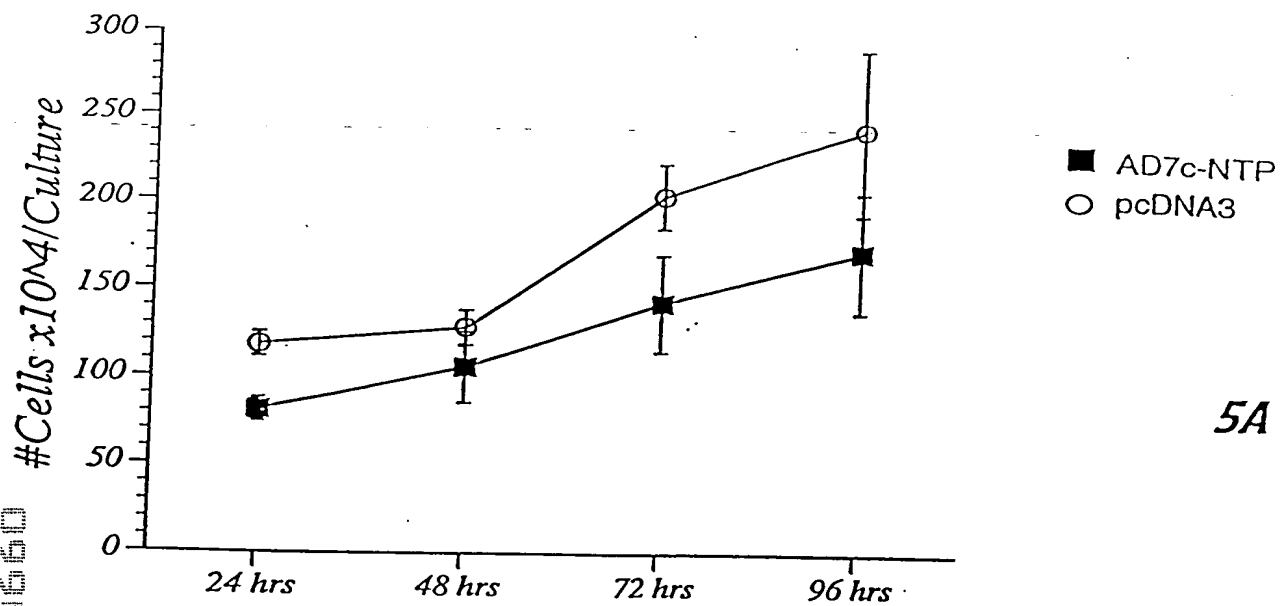


*FIGS. 3C-3F*

FIGS. 4A-4H

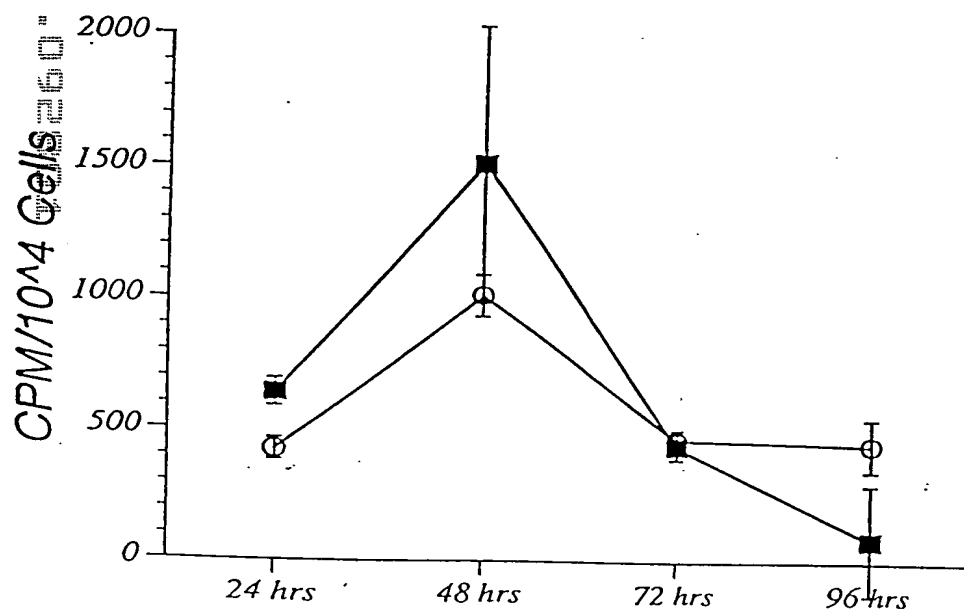


### Growth of SH-Sy5y Cells



5A

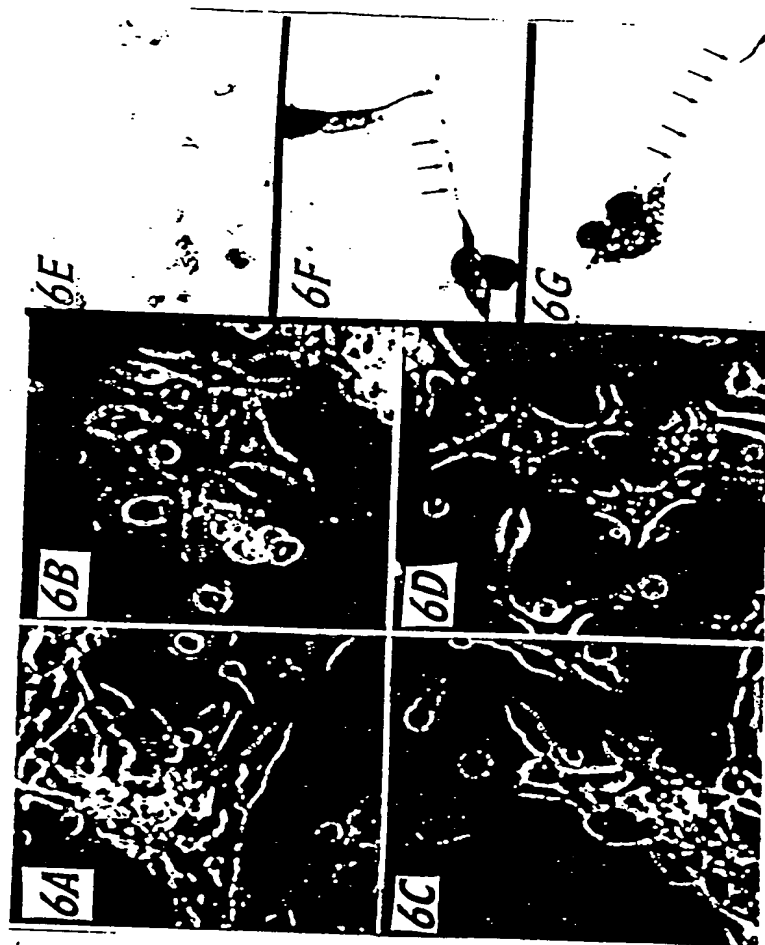
### <sup>3</sup>H-Thymidine Incorporation

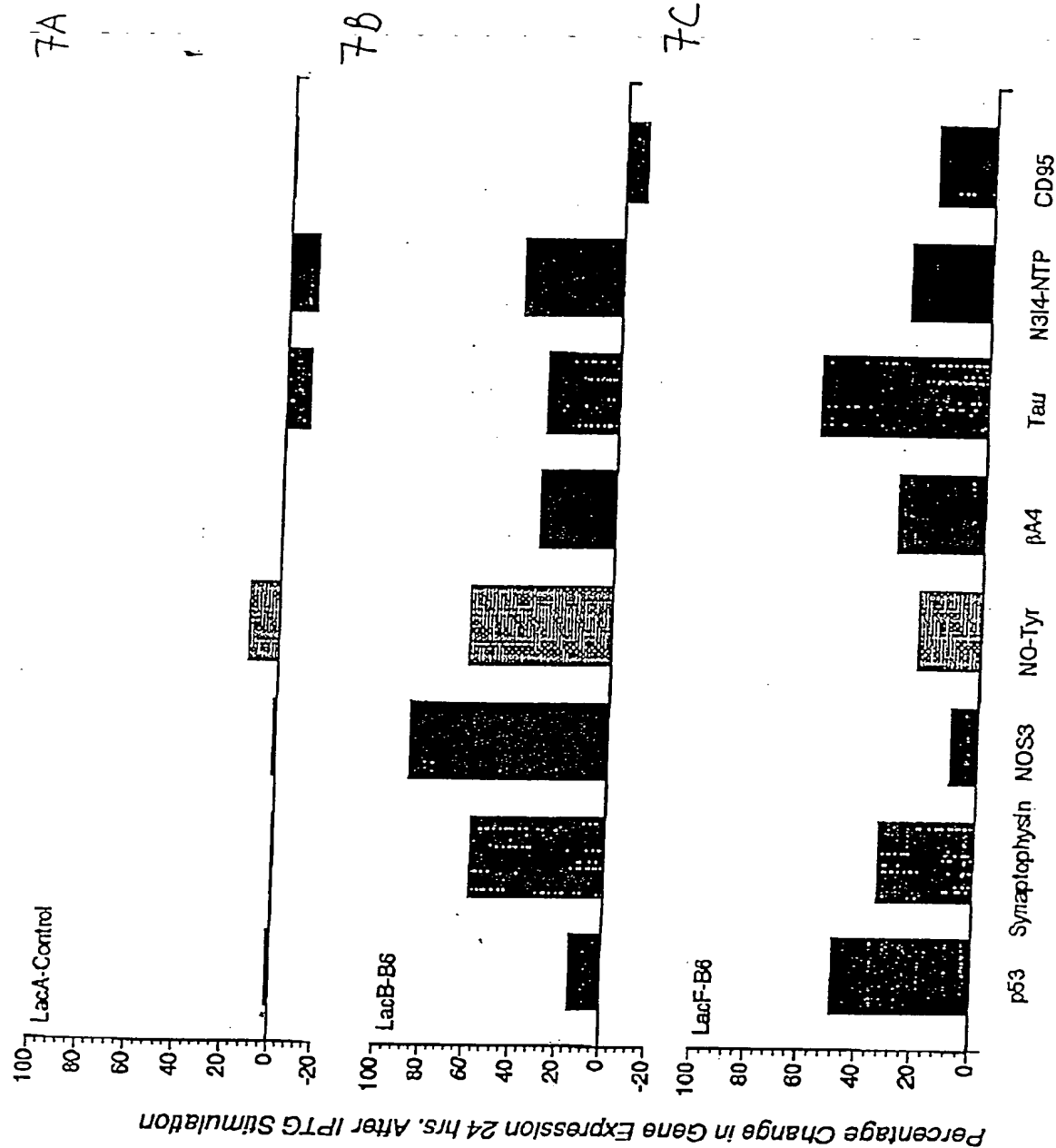


5B

FIGS. 5A-5B

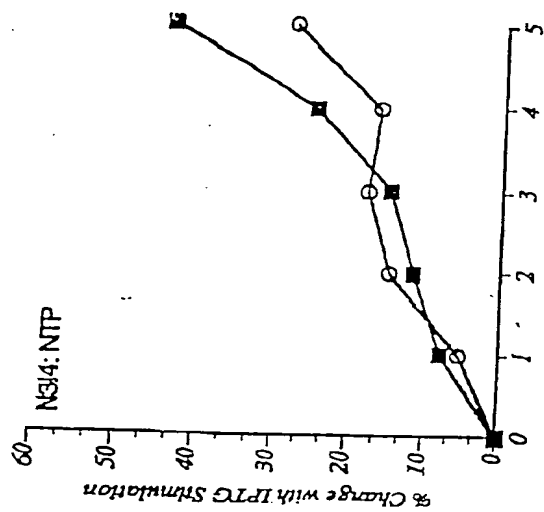
FIGS. 6A-6G



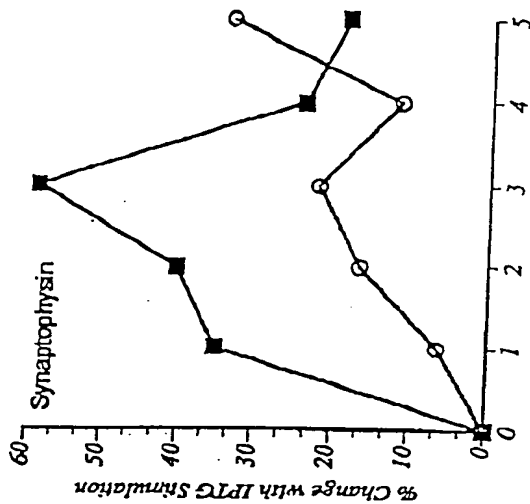




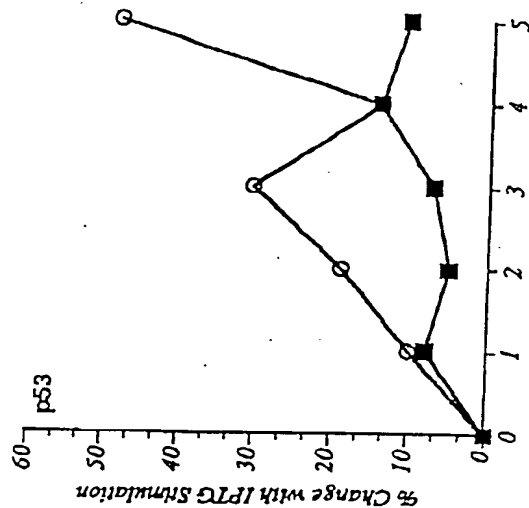
8A



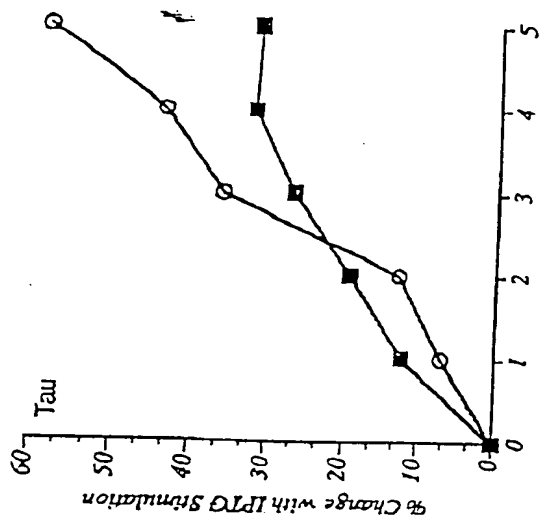
8B

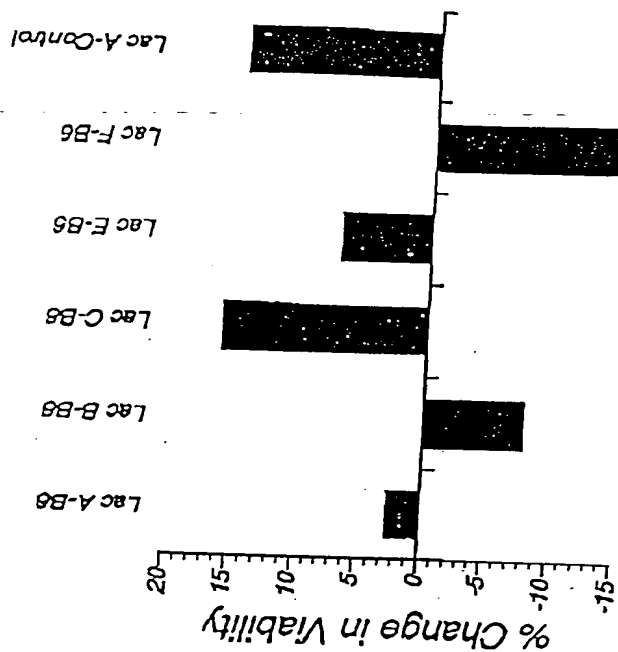
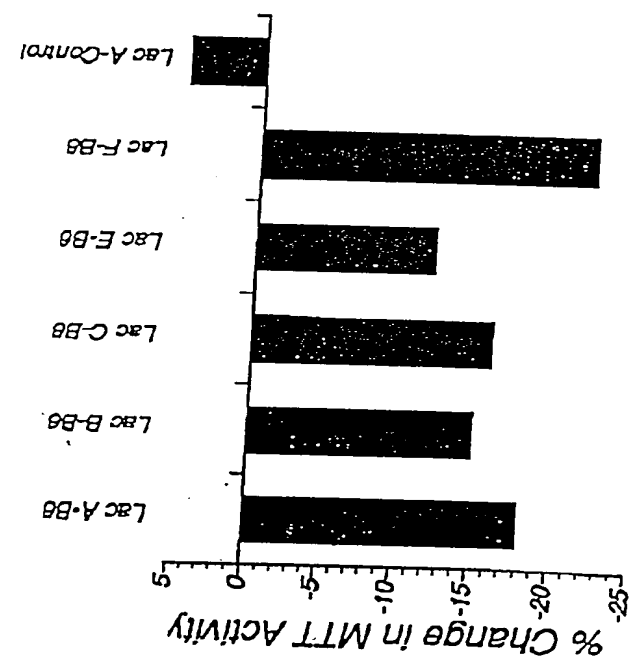


8D



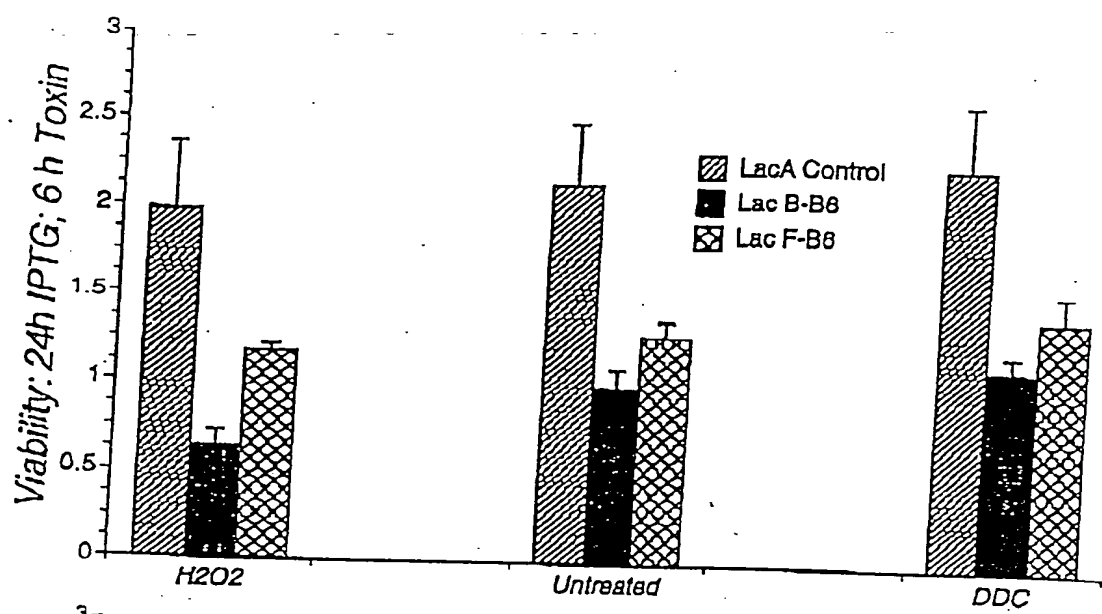
8C



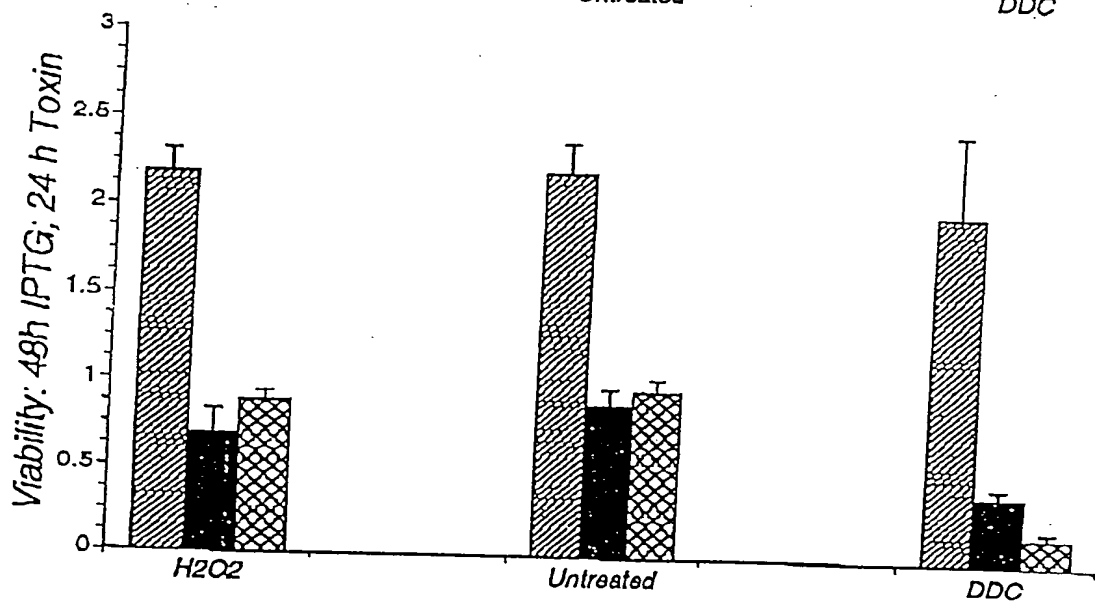


FO8260 109949660

10A



10B



FIGS. 10A-10B

FIG. 11

